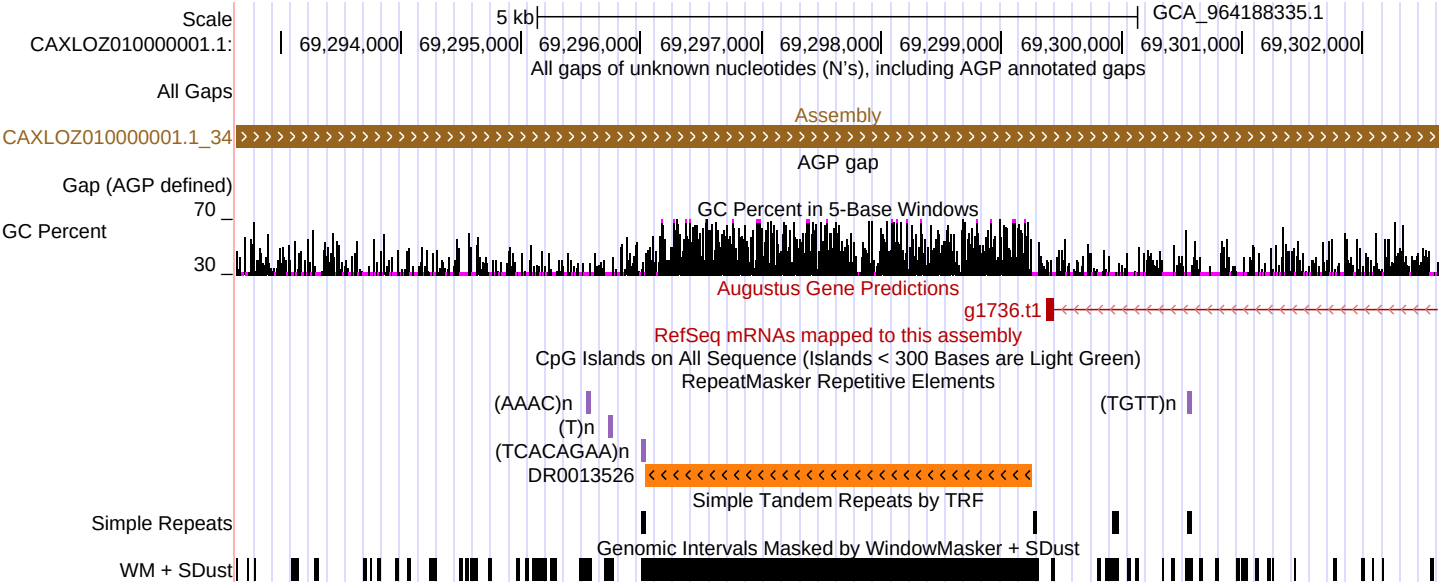
Scale
5 kb
GCA_964188335.1
CAXLOZ010000001.1:
69,294,000
69,295,000
69,296,000
69,297,000
69,298,000
69,299,000
69,300,000
69,301,000
69,302,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
CAXLOZ010000001.1_34
AGP gap
Gap (AGP defined)
70
GC Percent
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
g1736.t1
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AAAC)n
(TGTT)n
(T)n
(TCACAGAA)n
DR0013526
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust